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										E	DI-V	07 AV-	
252 :	D NO: 6								g	aat	RI ₇ Ki tcc Ser	acc'	0
atg gta a Met Val S SEQ I	ngc gct Ser Ala	att gt Ile Va	t tta l Leu	tat Tyr	gtg Val	ctt Leu	ttg Leu	gcg Ala	gcg Ala	gcg Ala	gcg Ala	cat His	48
64F1/ SEQ ID NO: 1													
tct gcc t	tt gcg	gcg ga	g cac	tgc	aac	gcg	caa	atg	aag	acg	ggt	ccg	96
Ser Ala P	Phe Ala	Ala Gl	u His	Cys	Asn	Ala	Gln	Met	Lys	Thr	Gly	Pro	
tac aag a	itt aaa	aac tt	g gac	att	acc	ccg	ccc	aag	gaa	acg	ctg	caa	144
Tyr Lys I	le Lys	Asn Le	u Asp	Ile	Thr	Pro	Pro	Lys	Glu	Thr	Leu	Gln	
aag gac g	gtg gaa	atc ac	c atc	gtg	gag	acg	gac	tac	aac	gaa	aac	gtg	192
Lys Asp V	/al Glu	Ile Th	r Ile	Val	Glu	Thr	Asp	Tyr	Asn	Glu	Asn	Val	
att atc g	gc tac Gly Tyr	aag gg Lys Gl	g tac y Tyr	tac Tyr	cag Gln	gcg Ala	tat Tyr	gcg Ala	tac Tyr	aac Asn	ggc Gly	ggc Gly	240
tcg ctg g	sat ccc	aac ac	a cgc	gtc	gaa	gaa	acc	atg	aaa	acg	ctg	aat	288
Ser Leu A	Isp Pro	Asn Th	r Arg	Val	Glu	Glu	Thr	Met	Lys	Thr	Leu	Asn	
gtg ggc a	aaa gag	gat tt	g ctt	atg	tgg	agc	atc	agg	cag	cag	tgc	gag	336
Val Gly L	.ys Glu	Asp Le	u Leu	Met	Trp	Ser	Ile	Arg	Gln	Gln	Cys	Glu	
gtg ggc g	gaa gag	ctg at	c gac	cgt	tgg	ggc	agt	gac	agc	gac	gac	tgt	384
Val Gly G	Glu Glu	Leu Il	e Asp	Arg	Trp	Gly	Ser	Asp	Ser	Asp	Asp	Cys	
ttt cgc g	sac aac	gag gg	c cgc	ggc	cag	tgg	gtc	aaa	ggc	aaa	gag	ttg	432
Phe Arg A	Asp Asn	Glu Gl	y Arg	Gly	Gln	Trp	Val	Lys	Gly	Lys	Glu	Leu	
gtg aag c	gg cag	aat aa	c aat	cac	ttt	gcg	cac	cac	acg	tgc	aac	aaa	480
Val Lys A	Irg Gln	Asn As	n Asn	His	Phe	Ala	His	His	Thr	Cys	Asn	Lys	
tcg tgg c	ga tgc	ggc 'at	t tcc	ac t	tcg	aaa	atg	tac	agc	agg	ctc	gag	528
Ser Trp A	Arg Cys	Gly Il	e Ser	Thr	Ser	Lys	Met	Tyr	Ser	Arg	Leu	Glu	
tgc cag g	sac gac	acg ga	c gag	tgc	cag	gta	tac	att	ttg	gac	gct	gag	576
Cys Gln A	Asp Asp	Thr As	o Glu	Cys	Gln	Val	Tyr	Ile	Leu	Asp	Ala	Glu	
ggc aac c Gly Asn P	cc atc	aac gt. Asn Va	acc Thr	gtg Val	gac Asp	act Thr	gtg Val	ctt Leu	cat His	cga Arg	gac Asp	ggc Gly	624
gtg agt a	itg att	ctc aa	a caa	aag	tct	acg	ttc	acc	acg	cgc	caa	ata	672
Val Ser M	let Ile	Leu Ly	s Gln	Lys	Ser	Thr	Phe	Thr	Thr	Arg	Gln	Ile	
aaa gct g Lys Ala A													720
gtg aca c	gc gaa	cac tg	t ttg	att	gac	aat	gat	ata	tat	gat	ctt	tct	768
Val Thr A	rg Glu	His Cy	Leu	Ile	Asp	Asn	Asp	Ile	Tyr	Asp	Leu	Ser	
aaa aac a	cg tgg	aac tge	aag	ttt	aac	aga	tgc	att	aaa	cgc	aaa	gtc	816
Lys Asn T	hr Trp	Asn Cy	Lys	Phe	Asn	Arg	Cys	Ile	Lys	Arg	Lys	Val	

FIG. 1